## Abstract

An isolated polynucleotide comprising a polynucleotide sequence selected from the group consisting of

- a) polynucleotide which is at least 70% identical to a
  polynucleotide that codes for a polypeptide which comprises the amino acid sequence of SEQ ID No. 2,
  - b) polynucleotide which codes for a polypeptide that comprises an amino acid sequence which is at least 70% identical to the amino acid sequence of SEQ ID No. 2,
- 10 c) polynucleotide which is complementary to the polynucleotides of a) or b), and
  - d) polynucleotide comprising at least 15 successive nucleotides of the polynucleotide sequence of a), b) or c),
- and processes for the fermentative preparation of L-amino acids using coryneform bacteria in which at least the metH gene is present in enhanced form, and use of the polynucleotide sequences as hybridization probes.

I:\atty\rwh\211714US.PA.doc